



US Seq List.ST25.txt  
SEQUENCE LISTING

RECEIVED

NOV 19 2002

TECH CENTER 1600/2900

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<120> Selection System

*Sub*  
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<140> 09/710,444

<141> 2000-11-20

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<151> 1998-05-13

<150> GB 9810228.8

<151> 1998-05-13

<150> PCT/GB99/01526

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<170> PatentIn version 3.1

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1 5 10 15

Glu

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<222> (1)..(89)

<223> Synthetic PCR primer for vector construction/screening

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 ccctcagaaa ggccggctgg gccgccacc 89

<210> 5

<211> 24

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gcgatggttg ttgtcattgt cggc

24

<210> 6

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<222> (1)..(24)

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aaaagaaacg caaagacacc acgg

24

<210> 7

<211> 23

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<210> 8

<211> 24

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<223> Synthetic PCR primer used to screen for recombinant clones.

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<222> (1)..(24)

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<210> 9

<211> 26

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 agcaa 65

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 <223> Synthetic PCT primer used to change codon usage in recombinant clones

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US Seq List.ST25.txt

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<212> DNA

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<223> Synthetic PCT primer used to change codon usage in recombinant clones

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gaaggaggtg gggtacccgg ttccgagggt gggtccggtt ccggtgattt tg

52

<210> 13

<211> 36

<212> DNA

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<222> (1)..(36)

<223> Synthetic PCR primer for vector construction/screening

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36

<210> 14

<211> 47

<212> DNA

<213> Bacillus amyloliquefaciens

<400> 14  
ctggcggcgg ccagccggc cctgcacagg ttatcaacac gtttgac

47

<210> 15

<211> 43

<212> DNA

<213> Bacillus amyloliquefaciens

<400> 15  
ctcggaaccg gtacctctga tttttgtaaa ggtctgataa gcg

43

<210> 16

<211> 44

<212> DNA

<213> Gallus gallus

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ggcggccag ccggcctttc tctctctgac gaggacttca aggc

44

<210> 17

<211> 41

<212> DNA

<213> Gallus gallus

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cctcggaacc ggtaccgaag agtcctttct ccttcttgag g

41

<210> 18

<211> 18

<212> DNA

<213> Artificial sequence

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<220>

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<222> (1)..(18)

<223> Synthetic PCR primer used for library construction

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tacgccaagc ttgcatgc

18

<210> 19

<211> 17

<212> DNA

<213> Artificial sequence

<220>

<223> Synthetic PCR primer used for library construction.

<220>

<221> misc\_feature

<222> (1)..(17)

<223> Synthetic PCR primer used for library constuction

<400> 19

ctgcacctgg gccatgg

17

<210> 20

<211> 17

<212> DNA

<213> Artificial sequence

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<220>

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<222> (1)..(17)

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<400> 20  
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17

<210> 21

<211> 126

<212> DNA

<213> *Erwinia chrysanthemi*

<220>

<221> misc\_feature

<223> n at positions 23, 24, 29, 55, 56, 81, 97, 101, and 102 can be G,  
A, T or C

<220>

<221> misc\_feature

<222> (23)..(23)

<223> n at position 23 can be G, A, T or C

<220>

<221> misc\_feature

<222> (24)..(24)

<223> n at position 24 can be G, A, T or C

<220>

<221> misc\_feature

<222> (29)..(29)

<223> n at position 29 can be G, A, T or C

<220>

<221> misc\_feature

<222> (55)..(55)

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<222> (56)..(56)

<223> n at position 56 can be G, A, T or C

<220>

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<222> (81)..(81)

<223> n at position 81 can be G, A, T or C

<220>

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<222> (97)..(97)

<223> n at position 97 can be G, A, T or C

<220>

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<222> (101)..(101)

<223> n at position 101 can be G, A, T or C

<220>

<221> misc\_feature

<222> (102)..(102)

<223> n at position 102 can be G, A, T or C

<400> 21

gattacgcca agcttgcatg canddctnt dtcaaggaga cagtcataat garrnrbcta 60

ttgsyaayrs yasyasyagb nttgttatta ctcsyanycv nncygdccat ggcccaggtg 120

cagctg 126

<210> 22

<211> 117

<212> DNA

<213> Bacteriophage M13mp18

<220>

<221> misc\_feature

<222> (18)..(18)

<223> Nucleotide at position 18 can be G, A, T or C.

<220>

<221> misc\_feature

<222> (19)..(19)

<223> Nucleotide at position 19 can be G, A, T or C.

<220>

<221> misc\_feature

<222> (20)..(20)

<223> Nucleotide at position 20 can be G, A, T or C.

<220>

<221> misc\_feature

<222> (21)..(21)

<223> Nucleotide at position 21 can be G, A, T or C.

<400> 22

gattacgcc a gctttg nnn ncttttttww ggagattttc aacrtgaraa rattattatt 60

csyaattsy ttagttsyts ytttctwtgy ggyccagccg gccatggccc aggtgca 117

<210> 23

<211> 18

<212> DNA

<213> Artificial sequence

<220>

<223> Synthetic PCR primer used for vector construction.

<400> 23

ctttatgctt ccggctcg

18

<210> 24

<211> 17

<212> DNA

<213> Artificial sequence

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<223> Synthetic PCR primer for library construction.

<220>

<221> misc\_feature

<222> (1)..(17)

<223> Synthetic PCT primer for library construction

<400> 24

cgccccatt cagatcc

17

<210> 25

<211> 50

<212> DNA

<213> Artificial sequence

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<223> Randomized E. chrysanthemi pelB sequence.

<220>

<221> misc\_feature

<222> (1)..(50)

<223> Randomized E. chrysanthemi pelB sequence

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aagcttgcat gcaaattcta tdtcaaggag acagttataa tgaaatacct

<210> 26

<211> 50

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<213> Artificial sequence

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<222> (1)..(50)

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B2 <222> (14)..(14)

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<223> n at position 15 can be G, A, T or C.

<220>

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<222> (20)..(20)

<223> n at position 20 can be G, A, T or C.

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<222> (45)..(45)

<223> n at position 45 can be G, A, T or C.

<220>

<221> misc\_feature

<222> (46)..(46)

<223> n at position 46 can be G, A, T or C.

<400> 26

aagcttgcat gcannddctn tdtcaaggag acagtcataa tgarrnbct

50

<210> 27

<211> 50

<212> DNA

<213> Artificial sequence

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B2

<220>

<221> misc\_feature

<222> (1)..(50)

<223> Randomized E. chrysanthemi pelB sequence

<400> 27

aagcttgcat gcagcatctc tdgcaaggag acagtcataa tgaagacgct

50

<210> 28

<211> 50

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<220>

<221> misc\_feature

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<222> (1)..(50)

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<400> 28

aagcttgcat gcacgggctg tdtcaaggag acagtcataa tgagagggct

50

<210> 29

<211> 50

<212> DNA

<213> Artificial sequence

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<221> misc\_feature

<222> (1)..(50)

<223> Randomized E. chrysanthemi pelB sequence

<400> 29

aagcttgcat gcaccagctc tdtcaaggag acagtcataa tgaggcggct

50

<210> 30

<211> 55

<212> DNA

<213> Artificial sequence

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<220>

<221> misc\_feature

<222> (1)..(55)

<223> Randomized E. chrysanthemi pelB sequence



<400> 30  
attcctaacg gcagccgctg gattgttatt actcgcggcc cagccggcca tggcc

55

<210> 31

<211> 55

<212> DNA

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<220>

<221> misc\_feature

<222> (1)..(55)

<223> Randomized E. chrysanthemi pelB sequence

<220>

<221> misc\_feature

<222> (22)..(22)

<223> n at position 22 can be G, A, T or C.

<220>

<221> misc\_feature

<222> (38)..(38)

<223> n at position 38 can be G, A, T or C.

<220>

<221> misc\_feature

<222> (42)..(42)

<223> n at position 42 can be G, A, T or C.

<220>

<221> misc\_feature

<222> (43)..(43)

<223> n at position 43 can be G, A, T or C.

<400> 31  
attgsyaayr syasyasyag bnttggttatt actcsyanyc vnncygdcca tggcc 55

<210> 32

<211> 55

<212> DNA

<213> Artificial sequence

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<220>

<221> misc\_feature

<222> (1)..(55)

<223> Randomized E. chrysanthemi pelB sequence

<400> 32  
attgcyaatg gtactgtyag gattggttatt actcccaccc ggtccgtcca tggcc 55

<210> 33

<211> 55

<212> DNA

<213> Artificial sequence

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<223> Randomized E. chrysanthemi pelB sequence.

<220>

<221> misc\_feature

<222> (1)..(55)

<223> Randomized E. chrysanthemi pelB sequence

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<210> 34

<211> 54

<212> DNA

<213> Artificial sequence

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<220>

<221> misc\_feature

<222> (1)..(54)

<223> Randomized E. chrysanthemi pe1B sequence

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<221> misc\_feature

<222> (22)..(22)

B2 <223> n at position 22 can be G, A, T or C.

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<221> misc\_feature

<222> (43)..(43)

<223> n at position 43 can be G, A, T or C.

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<221> misc\_feature

<222> (44)..(44)

<223> n at position 44 can be G, A, T or C.

<400> 34

attggtaata gcagcagtag bnttgtagg actcgacccc ccncyadcc atgg

54

<210> 35

<211> 22

US Seq List.ST25.txt

<212> PRT

<213> Erwinia chrysanthemi

<400> 35

Met Lys Tyr Leu Leu Pro Thr Ala Ala Ala Gly Leu Leu Leu Leu Ala  
1 5 10 15

Ala Gln Pro Ala Met Ala  
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<210> 36

<211> 22

<212> PRT

<213> Artificial sequence

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<221> MISC\_FEATURE

<222> (1)..(22)

<223> Randomized E. chrysanthemi pelB sequence

<400> 36

Met Lys Thr Leu Ala Met Val Leu Val Gly Gly Pro Pro Gly Pro Ser  
1 5 10 15

Ala Gln Pro Ala Met Ala  
20

<210> 37

<211> 21

<212> PRT

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<222> (1)..(21)

<223> Randomized E. chrysanthemi pelB sequence

<400> 37

Met Arg Gly Leu Ala Met Leu Val Ala Gly Gly Pro Ile Ala Pro Ala  
1 5 10 15

Gln Pro Ala Met Ala  
20

<210> 38

<211> 23

<212> PRT

<213> Artificial sequence

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<220>

<221> MISC\_FEATURE

<222> (1)..(23)

<223> Randomized E. chrysanthemi pelB sequence

<400> 38

Met Arg Arg Leu Val Pro Ile Thr Ala Ala Val Gly Leu Leu Ala Pro  
1 5 10 15

Pro Thr Gln Pro Ala Met Ala  
20

<210> 39

<211> 50

<212> DNA

<213> Artificial sequence

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<223> Randomized bacteriophage M13 g3 sequence.

<220>

<221> misc\_feature

<222> (1)..(50)

<223> Randomized bacteriophage M13 g3 sequence

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<210> 40

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<222> (9)..(9)

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<222> (1)..(50)

<223> Randomized bacteriophage M13 g3 sequence

<220>

<221> misc\_feature

<222> (10)..(10)

<223> n at position 10 is can be G, A, t or C.

<220>

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<222> (11)..(11)

<223> n at position 11 is can be G, A, t or C.

<220>

<221> misc\_feature

<222> (12)..(12)

<223> n at position 12 is can be G, A, t or C.

<400> 40

aagctttggn nncttttttw wggagatttt caacrtgara arattattat

50

<210> 41

<211> 50

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<213> Artificial sequence

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<220>

<221> misc\_feature

<222> (1)..(50)

<223> Randomized bacteriophage M13 g3 sequence.

<400> 41

aagctttggg gccttttttt aggagatttt caacatgaga agattattat

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<210> 42

<211> 50

<212> DNA

<213> Artificial sequence

US Seq List.ST25.txt

<220>

<223> Randomized bacteriophage M13 g3 sequence.

<220>

<221> misc\_feature

<222> (1)..(50)

<223> Randomized bacteriophage M13 g3 sequence

<400> 42

tcgcaattcc tttagttggt cttttctatg cggcccagcc ggccatggcc

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<210> 43

<211> 50

<212> DNA

<213> Artificial sequence

<220>

<223> Randomized bacteriophage M13 g3 sequence.

<220>

<221> misc\_feature

<222> (1)..(50)

<223> Randomized bacteriophage M13 g3 sequence

<400> 43

tcsyaattsy tttagttstt sytttctwtg yggycagcc ggccatggcc

50

<210> 44

<211> 50

<212> DNA

<213> Artificial sequence

<220>

<223> Randomized bacteriophage M13 g3 sequence.

<220>



&lt;221&gt; misc\_feature

&lt;222&gt; (1)..(50)

&lt;223&gt; Randomized bacteriophage M13 g3 sequence

&lt;400&gt; 44

tcctaattcc tttagtgtt gctttctatg tggccagcc ggccatggcc

50

&lt;210&gt; 45

&lt;211&gt; 22

&lt;212&gt; PRT

&lt;213&gt; Artificial sequence

&lt;220&gt;

&lt;223&gt; Randomized bacteriophage M13 g3 sequence.

&lt;220&gt;

&lt;221&gt; MISC\_FEATURE

&lt;222&gt; (1)..(22)

&lt;223&gt; Randomized bacteriophage M13 g3 sequence

&lt;400&gt; 45

Met Lys Lys Leu Leu Phe Ala Ile Pro Leu Val Val Pro Phe Tyr Ala  
 1 5 10 15

Ala Gln Pro Ala Met Ala  
 20

&lt;210&gt; 46

&lt;211&gt; 22

&lt;212&gt; PRT

&lt;213&gt; Artificial sequence

&lt;220&gt;

&lt;223&gt; Randomized bacteriophage M13 g3 sequence.

&lt;220&gt;

&lt;221&gt; MISC\_FEATURE

<222> (1)..(22)

<223> Randomized bacteriophage M13 g3 sequence

<400> 46

Met Arg Arg Leu Leu Leu Ala Pro Pro Val Ala Val Pro Phe Tyr Val  
1 5 10 15

Val Gln Pro Ala Met Ala  
20

<210> 47

<211> 18

<212> DNA

<213> Artificial sequence

<220>

<223> Synthetic oligonucleotide primer used as a substrate for Stoffel  
fragment of Thermus aquaticus DNA polymerase I.

32 <220>

<221> misc\_feature

<222> (1)..(18)

<223> Synthetic oligonucleotide primer used as substrate for Stoffel fr  
agment of Thermus aquaticus DNA polymerase I

<400> 47

tttcgcaaga tgtggcgt

18

<210> 48

<211> 12

<212> DNA

<213> Artificial sequence

<220>

<223> Synthetic oligonucleotide primer used as a substrate for Thermus  
aquaticus DNA polymerase I.

<220>

<221> misc\_feature

<222> (1)..(12)

<223> Synthetic primer used as substrate for Stoffel fragment of Thermus aquaticus DNA polymerase I

<400> 48

gcgaagatgt gg

12

<210> 49

<211> 30

<212> DNA

<213> Artificial sequence

<220>

<223> Synthetic oligonucleotide primer used as a substrate for Thermus aquaticus DNA polymerase I.

<220>

<221> misc\_feature

<222> (1)..(30)

<223> Synthetic oligonucleotide primer used as substrate for Thermus aquaticus DNA polymerase I

<400> 49

aaatacaaca ataaaacgcc acatcttgcg

30

<210> 50

<211> 20

<212> DNA

<213> Artificial sequence

<220>

<223> Synthetic oligonucleotide sequence insert containing PstI restriction site and frame shift for H102A mutant barnase fusion construct fused to p3 gene of phage fd-3.

<220>

<221> misc\_feature

<222> (1)..(20)

<223> Synthetic oligonucleotide sequence insert containing PstI restriction site and frame shift for H102A mutant barnase fusion construct fused to p3 gene of phage fd-3.

<400> 50  
ctgcaggcgg tgcggccgca

20

<210> 51

<211> 24

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<213> Artificial sequence

<220>

<223> Synthetic oligonucleotide used for random priming.

<220>

<221> misc\_feature

<222> (1)..(24)

B2 <223> Synthetic oligonucleotide used for random priming

<220>

<221> misc\_feature

<222> (19)..(19)

<223> n at position 19 can be G, A, T or C.

<220>

<221> misc\_feature

<222> (20)..(20)

<223> n at position 20 can be G, A, T or C.

<220>

<221> misc\_feature

<222> (21)..(21)

<223> n at position 21 can be G, A, T or C.

<220>

<221> misc\_feature

<222> (22)..(22)

<223> n at position 22 can be G, A, T or C.

<220>

<221> misc\_feature

<222> (23)..(23)

<223> n at position 23 can be G, A, T or C.

<220>

<221> misc\_feature

<222> (24)..(24)

<223> n at position 24 can be G, A, T or C.

B2

<400> 51  
gagcctgcag agctcaggnn nnnn

24

<210> 52

<211> 23

<212> DNA

<213> Artificial sequence

<220>

<223> Synthetic PCR primer used to re-amplify randomly amplified E. coli genomic DNA sequence.

<220>

<221> misc\_feature

<222> (1)..(23)

<223> Synthetic PCR primer used to re-amplify randomly amplified E. coli genomic DNA sequences.

<400> 52  
cgtgcgagcc tgcagagctc agg

23

<210> 53

<211> 45

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(45)

<223> Barstar binding barnase-p3 fusion insert

B2 <400> 53

Leu Gln Ser Ser Gly Asp Cys Val Ile Ser Asp Thr Cys Ile Ala Gly  
1 5 10 15

Met Ala Glu Ala Ala Ala Cys Glu Glu Lys Phe Ser Ser Gln Asn Val  
20 25 30

Gly Leu Thr Ile Thr Val Thr Pro Cys Leu Ser Ser Ala  
35 40 45

<210> 54

<211> 44

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(44)

<223> Barstar binding barnase-p3 fusion insert

<400> 54

Leu Gln Ser Ser Gly Cys Gly Ser Ser Gly Ser Ser Ile Asn Cys Leu  
1 5 10 15

Pro Cys Gly Ala Thr Ser Arg Gly Thr Ser Pro Leu Ala Ser Gly Leu  
20 25 30

Pro Ser Ser Ala Thr Ile His Cys Leu Ser Ser Ala  
35 40

<210> 55

<211> 40

<212> PRT

<213> Artificial sequence

B2 <220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(40)

<223> Barstar binding barnase-p3 fusion insert

<400> 55

Leu Gln Ser Ser Gly Asp Ser Ala Gly Cys Lys Asn Met Thr Gly Gly  
1 5 10 15

Arg Leu Tyr Ala His Thr Leu Glu Ala Ile Ile Pro Gly Phe Ala Val  
20 25 30

Ser Ala Pro Ala Cys Glu Pro Ala  
35 40

<210> 56

<211> 33

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(33)

<223> Barstar binding barnase-p3 fusion insert

<400> 56

Leu Gln Ser Ser Gly Cys Val Arg Leu Lys Arg Thr Ser Val Asn His  
1 5 10 15

Gln Pro Asp Ala Trp Pro Glu Pro His Leu Lys Ala Ala Cys Glu Pro  
20 25 30

Ala

<210> 57

<211> 44

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(44)

<223> Barstar binding barnase-p3 fusion insert

<400> 57

Leu Gln Ser Ser Gly Cys Gly Ser Ser Gly Ser Ser Ile Asn Cys Leu  
1 5 10 15

Pro Cys Gly Ala Thr Ser Arg Gly Thr Ser Pro Leu Ala Ser Gly Leu  
20 25 30



US Seq List.ST25.txt

Pro Ser Ser Ala Thr Val Gln Cys Leu Ser Ser Ala  
35 40

<210> 58

<211> 41

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(41)

<223> Barstar binding barnase-p3 fusion insert

B2 <400> 58

Leu Gln Ser Ser Gly Lys Ile Val Gln Ala Gly Ala Asn Ile Gln Asp  
1 5 10 15

Gly Cys Ile Met His Gly Tyr Cys Asp Thr Asp Thr Ile Val Gly Glu  
20 25 30

Asn Gly His Ile Gly Leu Ser Ser Ala  
35 40

<210> 59

<211> 45

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(45)

<223> Barstar binding barnase-p3 fusion insert

<400> 59

Leu Gln Ser Ser Gly Val Cys Val Ile Ser Asp Thr Cys Ile Ala Gly  
1 5 10 15

Thr Ala Glu Ala Ala Ala Cys Glu Glu Lys Phe Ser Ser Gln Asn Val  
20 25 30

Gly His Thr Ile Thr Glu Thr Pro Cys Leu Ser Ser Ala  
35 40 45

<210> 60

<211> 44

<212> PRT

<213> Artificial sequence

32 <220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(44)

<223> Barstar binding barnase-p3 fusion insert

<400> 60

Leu Gln Ser Ser Gly Cys Gly Ser Ser Gly Ser Ser Ile Asn Cys Leu  
1 5 10 15

Pro Cys Gly Ala Thr Ser Arg Gly Thr Ser Pro Leu Ala Ser Gly Leu  
20 25 30

Pro Ser Ser Ala Thr Ile Gln Cys Leu Ser Ser Ala  
35 40

<210> 61

<211> 53

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(53)

<223> Barstar binding barnase-p3 fusion insert

<400> 61

Leu Gln Ser Ser Gly Gln Asp Ser Gln Arg Glu His Ala Ser His Thr  
1 5 10 15

Ala Glu Asp Asp Cys Glu Asp Gln Thr Arg Ile His Gln His Ile Arg  
20 25 30

Glu Val Asp Phe Val Asp Thr Pro Gln Glu Val Asp Asp Cys Arg Ala  
35 40 45

B2 Ala Leu Ser Ser Ala  
50

<210> 62

<211> 33

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(33)

<223> Barstar binding barnase-p3 fusion insert

<400> 62

Leu Gln Ser Ser Gly Cys Val Arg Leu Lys Arg Thr Ser Val Asn His  
Page 35

1

5

10

15

Gln Pro Asp Ala Trp Pro Glu Pro His Leu Lys Ala Ala Cys Glu Pro  
20 25 30

Ala

&lt;210&gt; 63

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Artificial sequence

&lt;220&gt;

&lt;223&gt; Barstar binding barnase-p3 fusion insert.

&lt;220&gt;

&lt;221&gt; MISC\_FEATURE

&lt;222&gt; (1)..(9)

&lt;223&gt; Barstar binding barnase-p3 fusion insert

&lt;400&gt; 63

Leu Gln Ser Ser Gly Val Arg Pro Ala  
1 5

&lt;210&gt; 64

&lt;211&gt; 44

&lt;212&gt; PRT

&lt;213&gt; Artificial sequence

&lt;220&gt;

&lt;223&gt; Barstar binding barnase-p3 fusion insert.

&lt;220&gt;

&lt;221&gt; MISC\_FEATURE

&lt;222&gt; (1)..(44)

&lt;223&gt; Barstar binding barnase-p3 fusion insert

US Seq List.ST25.txt

<400> 64

Leu Gln Ser Ser Gly Cys Gly Ser Ser Gly Ser Ser Ile Asn Cys Leu  
1 5 10 15

Pro Cys Gly Ala Thr Ser Arg Gly Thr Ser Pro Leu Ala Ser Gly Leu  
20 25 30

Pro Ser Ser Ala Thr Ile Gln Cys Leu Ser Ser Ala  
35 40

<210> 65

<211> 30

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(30)

<223> Barstar binding barnase-p3 fusion insert

<400> 65

Leu Gln Ser Ser Gly Thr Glu Val Asp Arg Gly Asn Gln Gln His Asp  
1 5 10 15

Thr Asn Asp Arg Asp Phe Thr His Thr Pro Leu Ser Ser Ala  
20 25 30

<210> 66

<211> 36

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

US Seq List.ST25.txt

<220>

<221> MISC\_FEATURE

<222> (1)..(36)

<223> Barstar binding barnase-p3 fusion insert

<400> 66

Leu Gln Ser Ser Gly Val Ala Gln Gly Ser Ser Ala Ser Val Asp Val  
1 5 10 15

Thr Ala Thr Asn Ala Val Leu Ser Ala Asp Ser Leu Ser Leu Gly Gly  
20 25 30

Gly Glu Pro Ala  
35

<210> 67

<211> 19

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(19)

<223> Barstar binding barnase-p3 fusion insert

<400> 67

Leu Gln Ser Ser Gly Gly Ala Val Ala Val Thr Pro Gly Pro Val Leu  
1 5 10 15

Ser Ser Ala

<210> 68

<211> 18

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(18)

<223> Barstar binding barnase-p3 fusion insert

<400> 68

Leu Gln Ser Ser Gly His Cys Arg Gly Lys Pro Val Leu Cys Thr His  
1 5 10 15

Thr Ala

<210> 69

<211> 9

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(9)

<223> Barstar binding barnase-p3 fusion insert

<400> 69

Leu Gln Ser Ser Gly Val Arg Pro Ala  
1 5

<210> 70

<211> 36

US Seq List.ST25.txt

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(36)

<223> Barstar binding barnase-p3 fusion insert

<400> 70

Leu Gln Ser Ser Gly Glu Pro Ala Pro Ala His Glu Ala Lys Pro Thr  
1 5 10 15

Glu Ala Pro Val Ala Lys Ala Glu Ala Lys Pro Glu Thr Pro Ala His  
20 25 30

B2 Leu Ser Ser Ala  
35

<210> 71

<211> 33

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(33)

<223> Barstar binding barnase-p3 fusion insert

<400> 71

Leu Gln Ser Ser Gly Cys Val Arg Leu Lys Arg Thr Ser Val Asn His  
1 5 10 15



US Seq List.ST25.txt

Gln Pro Asp Ala Trp Pro Glu Pro His Leu Lys Ala Ala Cys Glu Pro  
20 25 30

Ala

<210> 72

<211> 36

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(36)

<223> Barstar binding barnase-p3 fusion insert

<400> 72

Leu Gln Ser Ser Gly Val Val Asp Trp Ala Lys Met Arg Glu Ile Ala  
1 5 10 15

Asp Ser Ile Gly Ala Tyr Leu Phe Val Asp Met Ala His Val Ala Ala  
20 25 30

Leu Ser Ser Ala  
35

<210> 73

<211> 117

<212> DNA

<213> Artificial sequence

<220>

<223> Vector pK1 polylinker sequence.

<220>

US Seq List.ST25.txt

<221> misc\_feature

<222> (1)..(117)

<223> Vector pK1 polylinker sequence

<400> 73

aatgctggcg gcggcccagc cggcctttct gaggggtcga ctatagaagg acgagggggcc 60

cacgaaggag gtgggggtacc cggttccgag ggtggttccg gttccggtga ttttgat 117

<210> 74

<211> 39

<212> PRT

<213> Artificial sequence

<220>

<223> Polypeptide encoded by pK1 vector polylinker sequence.

<220>

<221> MISC\_FEATURE

<222> (1)..(39)

<223> Polypeptide encoded by pK1 vector polylinker sequence

<400> 74

Asn Ala Gly Gly Gly Pro Ala Gly Leu Ser Glu Gly Ser Thr Ile Glu  
1 5 10 15

Gly Arg Gly Ala His Glu Gly Gly Gly Val Pro Gly Ser Glu Gly Gly  
20 25 30

Ser Gly Ser Gly Asp Phe Asp  
35

<210> 75

<211> 117

<212> DNA

<213> Artificial sequence

US Seq List.ST25.txt

<220>

<223> Vector pk2 polylinker sequence.

<220>

<221> misc\_feature

<222> (1)..(117)

<223> vector pk2 polylinker sequence

<400> 75

aatgctggcg gcggcccagc cggcctttct gaggggtcga ctatagaagg acgagggccc 60

acgaagcagc tgggggtaccg gttccgaggg tggttccggt tccggtgatt ttgatta 117

<210> 76

<211> 39

<212> PRT

<213> Artificial sequence

B2

<220>

<223> Polypeptide sequence encoded by vector pk2 polylinker region.

<220>

<221> MISC\_FEATURE

<222> (1)..(39)

<223> Polypeptide sequence encoded by vector pk2 polylinker region.

<220>

<221> MISC\_FEATURE

<222> (38)..(38)

<223> X represents a TGA stop codon

<220>

<221> MISC\_FEATURE

<222> (36)..(36)

<223> X represents a stop codon (TGA)

US Seq List.ST25.txt

<400> 76

Asn Ala Gly Gly Gly Pro Ala Gly Leu Ser Glu Gly Ser Thr Ile Glu  
1 5 10 15

Gly Arg Gly Pro Thr Lys Gln Leu Gly Tyr Arg Phe Arg Gly Trp Phe  
20 25 30

Arg Phe Arg Xaa Phe Xaa Leu  
35

<210> 77

<211> 35

<212> DNA

<213> Artificial sequence

<220>

<223> Sequence of the junction region between Barnase and p3 in recombi  
nant fusion vector fd-3.

<220>

<221> misc\_feature

<222> (1)..(35)

<223> Sequence of the junction region between Barnase and p3 in recombi  
nant fusion vector fd-3.

<400> 77

atcagactgc aggcggtgcg gccgcagaaa ctggt

35

<210> 78

<211> 11

<212> PRT

<213> Artificial sequence

<220>

<223> Amino acid sequence about the junction of Barnase and p3 coding r  
egions of recombinant fusion vector fd-3.

<400> 78

Ile Arg Leu Gln Ala Ala Ala Ala Glu Thr Val

1

5

<210> 79

<211> 4

<212> PRT

<213> Artificial sequence

*Sub*  
*optimal*  
<220>

<223> Factor Xa protease cleavage sequence.

<220>

<221> MISC\_FEATURE

<222> (1)..(1)

*B2*  
*cons*  
<223> X can be either Ile or Leu.

<220>

<221> MISC\_FEATURE

<222> (1)..(4)

<223> Factor Xa proteolytic cleavage site.

<400> 79

Xaa Glu Gly Arg  
1